# Figure 1-A

				mmaaaamaam	000100000
>g2804590	CCTGCCTGCA	CCCGC'I': CAC	CCTGAGCGCC	TTGGGGTGGT	TGCAGACTGT
>g2804590	CCMCCCMCCA	CCCCCM, CAC	CCTTCANCCCC		
>2682428 >2682469			-	TTGGGGTGGT	
	CCTGCCTGCA	CCCGCTTCAC	CCTGAGCGCC		TGCAGACTGT
>3359342				GGAGGC	
>1736231					TGCAGACTGT
>1734520				mmaaaamaam	TGCAGACTGT
Consensus	CCTGCCTGCA	CCCGCT: CAC	CCTGAGCGCC	TTGGGGTGGT	TGCAGACTGT
>g2804590	GGAATCCCCA	CTGTGCAG			
>g2804590	GGAGCCGGGA	GCCGGCAG			
>g2804590		CAGTA	AGCCCAGAGG	TCTCCACCCC	A:CGGGAGGA
>2682428	GGAATCCCCA	CTGTGCAGTA	AGCCCAGATG	TCTCCANNCC	AACGGTAGGA
>2682469	GGAATCCCCA	CTGTGCAGTA	AGCCCAGAGG	TCTCCACCCC	A:CGGGATGA
>3359342	GGAGCCGGGA	GCCGGCAGTA	AGCCCAGAGG	TCTCCACCCC	A:CGGGAGGA
>1736231	GGAGCCGGGA	GCCGGCAGTA	AGCCCAGAGG	TCTCCACCCC	A:CGGGAGGA
>1734520	GGAGCCGGGA	GCCGGCAGTA	AGCCCAGAGG	TCTCCACCCC	A: CGGGAGGA
Consensus	GGAGCCGGGA	GCCGGCAGTA	AGCCCAGAGG	TCTCCACCCC	A:CGGGAGGA
2004500	N GGGMGN GGG	CAACACCCCC	CAACACAMCC	ACCGCGTGAC	CA CAMA CCCC
>g2804590 >2682428				ANCGCGTGAC	
>2682428				ACCGCGTGAC	
>3359342				ACCGCGTGAC	
>1736231				ACCGCGTGAC	
>1736231				ACCGCGTGAC	
>2596108	AGGC TGAGGC			ACCGCGTGAC	
Consensus	AGGCTGAGGC			ACCGCGTGAC	
>g2804590	ATCCTGGGCA	TCCCTCAGGC	ACACCGTGG:	CACCGGCCTG	GTGCTGGATG
>2682428	ATCCTGNGCA	TCCCTCAGGC	ACACCGTGG:	CACCGGCCTG	GTGCTGGATG
>2682469	ATCCTGGGCA	TCCCTCAGGC	ACACCGTGGG	CACCGGCCTG	GTGCTNGATG
>3359342	ATCCTGGGCA	TCCCTCAGGC	ACACCGTGG:	CACCGGT	
>1736231	ATCCTGGGC				
>1734520	· · ·	TCCCTCAGGC	•		GTGCTGGATG
>2596108			ACACCGTGG:		GTGCTGGATG
Consensus	ATCCTGGGCA	TCCCTCAGGC	ACACCGTGG:	CACCGGCCTG	GTGCTGGATG
>g2804590	GAGACACCAG	CTACACATAC	CATCTGGTGT	GCATGGGCCC	CGAGGCCAGC
>2682428				GCATTGGCCC	
>2682469				GCATGGGCCC	
>1734520				GCATGGGCCC	
>2596108				GCATGGGCCC	
>3388863	0.101.01.001.0	01		30	TAGGGCCAGC
Consensus	GAGACACCAG	СТАСАСАТАС	CATCTGGTGT	GCATGGGCCC	CGAGGCCAGC
consensus	ononeneeno	cincincinin	cc.100101	Genrodece	conocence
>g2804590					ACAGGGCCCA
>2682428			GCA: ACATGG		
>2682469			GCAGACATGG	CC: ACTGACC	NAAGGCCAT
>1734520	GG: TGGGGCC				
>2596108					ACAGGGCCCA
>3388863					ACAGGGCCCA
Consensus	GGCTGGGGCC	AGGATGAGCC	GCAGACATGG	CCCACTGACC	ACAGGGCCCA

# Figure 1-B

>g2804590				CAGCGTGCAT	
>2596108	GCAGGGCGTG	CAGAGGCAGG	GGGTGTCCTA	CAGCGTGCAT	GCCTACACTG
>3388863				CAGCGTGCAT	
Consensus	GCAGGGCGTG	CAGAGGCAGG	GGGTGTCCTA	CAGCGTGCAT	GCCTACACTG
>g2804590				AGAACAGGGA	GGATGAGGGT
>2596108	GCCAGCCGTC				
>3388863				AGAACAGGGA	
Consensus	GCCAGCCGTC	CCCACGGGGG	CTCCACTCGG	AGAACAGGGA	GGATGAGGGT
>g2804590	TGGCAGGTTT	ACCGCCTGGG	CGCCAGGGAT	GCCCACCAGG	GACGTCCAAC
>3388863	TGGCAGGTTT	ACCGCCTGGG	CGCCAGGGAT	GCCCACCAGG	GACGTCCAAC
Consensus	TGGCAGGTTT	ACCGCCTGGG	CGCCAGGGAT	GCCCACCAGG	GACGTCCAAC
>g2804590	ATGGGCACTC	CGCCCAGAGG	ACGGGGAGGA	CAAGGAGATG	AAGACCTACC
>3388863	ATGGGCACTC	CGCCCAGAGG	ACGGGGAGGA	CAAGGAGATG	AAGACCTACC
<q2322685< td=""><td></td><td>GCCCAGAGG</td><td>ACGGGGAGGA</td><td>CAAGGAGATG</td><td>AAGACCTACC</td></q2322685<>		GCCCAGAGG	ACGGGGAGGA	CAAGGAGATG	AAGACCTACC
Consensus	ATGGGCACTC	CGCCCAGAGG	ACGGGGAGGA	CAAGGAGATG	AAGACCTACC
>g2804590				GGCTGTGTGA	
>3388863				GGCTGTGTGA	
<g2322685< td=""><td></td><td></td><td></td><td>GGCTGTGTGA</td><td></td></g2322685<>				GGCTGTGTGA	
Consensus	GCCTGGATGC	TGGGGACGCT	GACCCCAGGA	GGCTGTGTGA	CCTGGAGCGG
>g2804590	GAGCGCTGGG	CCGTCATCCA	GGGCCAGGCA	GTCAGGAAGA	GCAGCACCGT
<g2322685< td=""><td>GAGCGCTGGG</td><td>CCGTCATCCA</td><td>GGGCCAGGCA</td><td>GTCAGGAAGA</td><td>GCAGCACCGT</td></g2322685<>	GAGCGCTGGG	CCGTCATCCA	GGGCCAGGCA	GTCAGGAAGA	GCAGCACCGT
Consensus	GAGCGCTGGG	CCGTCATCCA	GGGCCAGGCA	GTCAGGAAGA	GCAGCACCGT
>g2804590				AGACCCCAGG	
<g2322685< td=""><td></td><td></td><td></td><td>AGACCCCAGG</td><td></td></g2322685<>				AGACCCCAGG	
Consensus	GGCCACGCTC	CAGGGCACTC	CTGACCACGG	AGACCCCAGG	ACCCCCGGCC
>g2804590	CACCTCGGTC	CACGCCCCTG	GAGGAGAACG	TGGTTGACAG	GGAGCAGATT
<g2322685< td=""><td>CACCTCGGTC</td><td>CACGCCCCTG</td><td>GAGGAGAACG</td><td>TGGTTGACAG</td><td>GGAGCAGATT</td></g2322685<>	CACCTCGGTC	CACGCCCCTG	GAGGAGAACG	TGGTTGACAG	GGAGCAGATT
Consensus	CACCTCGGTC	CACGCCCCTG	GAGGAGAACG	TGGTTGACAG	GGAGCAGATT
>g2804590				AGTCTGGAGC	AGGCGAACAA
<g2322685< td=""><td>GACTTCCTGG</td><td>CAGCGAGACA</td><td>GC</td><td></td><td></td></g2322685<>	GACTTCCTGG	CAGCGAGACA	GC		
>3988413					AGGCGAACAA
Consensus	GACTTCCTGG	CAGCGAGACA	GCAGTTCCTG	AGTCTGGAGC	AGGCGAACAA
>g2804590	GGGGGCCCCT	CATAGCTCCC	CGGCCAGGG	GACCCCTGCA	GGCACAACCC
>3988413	GGGGGCCCCT	CATAGCTCCC	CGGCCNGGGG	GACCCCTGCA	GGCACAACCC
Consensus	GGGGGCCCCT	CATAGCTCCC	CGGCCAGGGG	GACCCCTGCA	GGCACAACCC
>g2804590					CCTGGCCAAC
>3988413					CCTGGCCAAC
Consensus	CAGGGGCCAG	CCAGGCCCCC	AAGGCCTTCA	ACAAGCCCCA	CCTGGCCAAC
>g2804590	GGGCACGTGG	TTCCCATCA	GCCCCAGGTG	AAGGGGGTGG	TCAGGGAAGA
>3988413					TCAGGGAAGA
Consensus					TCAGGGAAGA

#### Figure 1-C

>q2804590	GAACAAGGTG	CGTGCTGTGC	CCACCTGGGC	CAGTGTCCAA	GTTGTGGATG
>3988413			CCACCTGGGC		
Consensus	GAACAAGGTG	CGTGCTGTGC	CCACCTGGGC	CAGTGTCCAA	GTTGTGGATG
-,					
>q2804590	ACCCTGGCTC	CTTGGCCTCA	GTGGAGTCCC	CGGGGACCCC	CAAGGAGACG
>3988413	ACCCTGGCTC	CTTGGCCTCA	GTGGAGTCCC	CGGGGACCCC	CAAGGAGACG
Consensus			GTGGAGTCCC		
>g2804590	CCCATCGAGC	GGGAGATCCG	TCTGGCTCAG	${\tt GAGCGTGAGG}$	CAGACCTGCG
>3988413	CCCATCGAGC	GGGAGATCCG	TCTGGCTCAG		
Consensus	CCCATCGAGC	GGGAGATCCG	TCTGGCTCAG	GAGCGTGAGG	CAGACCTGCG
•					
>g2804590			AGGCAACCGA		
Consensus	AGAGCAGAGG	GGGCTTCGGC	AGGCAACCGA	CCACCAGGAG	CTGGTGGAAA
		<u>.</u>			
>g2804590			ACCAAGCTGA		
Consensus	TCCCCACCAG	GCCGCTGCTG	ACCAAGCTGA	GCCTGATCAC	AGCCCCACGG
2004500	0000101010	accacama	CCTCTACGTG	030000-030	A #12 C#13 C A CC
>g2804590	CGGGAGAGAG	GGCGCCCGTC			
>3615515	0000101010	GGGGGGGGTG		CAGCGGNGAC	
Consensus	CGGGAGAGAG	GGCGCCCGTC	CCTCTACGTG	CAGCGGNGAC	ATAGTACAGG
>g2804590	ACACACACOC	TCACCAACAC	CACCGGCGGG	AGGGCCTGCA	CCTCCCCCC
>3615515			CACCGGCGGG		
Consensus			CACCGGCGGG		
Consensus	AGACACAGCG	TOROGRAGAC	CACCOGCOGO	nadace roen	0010000000
>g2804590	GCGTCCACAC	CCGACTGGGT	CTCGGAGGGT	CCCCAGCCCG	GACTCCGGAG
>3615515	GCGTCCACAC	CCGACTGGGT	CTCGGAGGGT	CCCCAGCCCG	GACTCCGGAG
Consensus	GCGTCCACAC	CCGACTGGGT	CTCGGAGGGT	CCCCAGCCCG	GACTCCGGAG
>g2804590	AGCCCTCAGC	TCAGATTCCA	TCCTCAGCCC	GGCCCCAGAT	GCCCGTGCGG
>3615515			TCCTCAGCCC		
Consensus	AGCCCTCAGC	TCAGATTCCA	TCCTCAGCCC	GGCCCCAGAT	GCCCGTGCGG
					•
>g2804590			AGGAAGGTGA		
>3615515	CCGACCCAGC	TCCAGAAGTG	AGGAAGGTGA	ACCGCATCCC	-
>2055371					CTGATGCC
>2055371IH					TGATGCC
Consensus	CCGACCCAGC	TCCAGAAGTG	AGGAAGGTGA	ACCGCATCCC	ACCTGATGCC
>g2804590	ma cca ccccm	እሮርሞርእርርርር	CCCCACCCCC	ር አርርጥአር አአጥ	TCTCAGCCTT
>3615515					TCTCAGC: TT
>2055371					TCTCAGCCTT
>2055371					TCTCAGCCTT
					TCTCAGCCTT
Consensus	TACCAGCCGT	ACCIGAGCCC	CGGGACCCCC	CAGCIAGAAI	TCTCAGCCTT
>g2804590	CGGAGCATTC	GGCAAGCCCA	GCAGTCTCTC	CACAGCGGAG	G:CCAAGGCT
>3615515		GGCAAGCCCA		•	
>2055371				NACAGCGGAG	GANCAAGGCT
>2055371IH					G:CCAAGGCT
>1431231			CTCTC	CACAGCGGAG	G:CCAAGGCT
Consensus	CGGAGCATTC	GGCAAGCCCA	GCAGTCTCTC	CACAGCGGAG	G:CCAAGGCT

# Figure 1-D

-2004500	CCCACMMCAC	CAAACCCCAC	CARCTCCCCC	AGGCATCTCT	CACAATICCTC
>g2804590 >2055371				AGGCATCTCT	
>2055371 >2055371IH				AGGCATCTCT	
>20333711H >1431231				AGGCATCTCT	
				AGGCATCTCT	
Consensus	GCGACTICAC	CAAAGGCCAC	GAIGICCCCG	AGGCATCTCT	CAGAATCCTC
>g2804590	TGGAAAACCC	CTGAGCACAA	AGCAAGAGGC	ATCGAAGCCC	CCTCGGGGAT
>2055371	TGGAAAACCC	CTGAGCACAA	AGCAAGAGGC	ATCGAAGCCC	CCTCGGGGAT
>2055371IH	TGGAAAACCC	CTGAGCACAA	AGCAAGAGGC	ATCGAAGCCC	CCTCGGGGAT
>1431231	TGGAAAACCC	CTGAGCACAA	AGCAAGAGGC	ATCGAAGCCC	CCTCGGGGAT
>3253860		•			CGGGGAT
Consensus	TGGAAAACCC	CTGAGCACAA	AGCAAGAGGC	ATCGAAGCCC	CCTCGGGGAT
>g2804590	GCCCGCAAGC	CAACAGGGGT	GTCGTGCGGT	GGGAGTACTT	CCGCCTGCGT
>2055371	GCCCGCAAGC	CAACAGGG			
>2055371IH	GCCCGCAAGC	CAACAGGGGT	GTCGTGCGGT	GGGAGTACTT	CCGCCTGCGT
>1431231	GCCCGCAAGC	CAACAGGGGT	GTCGTGCGGT	GGGAGTACTT	CCGCCTGCGT
>3253860				GGGAGTACTT	
Consensus				GGGAGTACTT	
	33333312133				0000010001
>g2804590	CCTCTGCGGT	TCAGGGCCCC	AGACGAGCCC	CAGCAGGCCC	AAGTCCCCCA
>2055371IH	CCTCTGCGGT	TCAGGGCCCC	AGACGAGCCC	CAGCAGGCCC	AAGTCCCCCA
>1431231	CCTCTGCGGT	TCAGGGCCCC	AGACGAGCCC	CA	
>3253860	CCTCTGCGGT	TCAGGGCCCC	AGACGANCCC	CAGCAGGCCC	AAGTCCCCCA
Consensus	CCTCTGCGGT	TCAGGGCCCC	AGACGAGCCC	CAGCAGGCCC	AAGTCCCCCA
>g2804590				TGCACTGAGG	
>2055371IH				TGCACTGAGG	
>3253860	TGTCTGGGGC			TGCACTGAGG	· · - · · · <del>-</del> -
>1753756				NGCACTGAGG	
Consensus	TGTCTGGGGC	TGGGAGGTGG	CTGGGGCCCC	TGCACTGAGG	CTGCAGAAGT
>g2804590	CCCAGTCATC	TGATCTGCTG	GAAAGGGAGA	GGGAGAGTGT	CCTGCGCCGG
>2055371IH	CCCAGTCATC	TGATCTGCTG	GAAAGGGAGA	GGGAGAGTGT	CCTGCGCCGG
>3253860	CCCAGTCATC	TGATCTGCTG	GAAAGGGAGA	GGGAGAGTGT	CCTGCGCCGG
>1753756	CCCAGTCATC	TGATCTGCTG	GAAAGGGAGA	GGGAGAGTGT	CCTGCGCCGG
Consensus	CCCAGTCATC	TGATCTGCTG	GAAAGGGAGA	GGGAGAGTGT	CCTGCGCCGG
>q2804590	GAGCAAGAGG	TGGCAGAGGA	GCGGAGAAAT	GCTCTCTTCC	CAGAGGTCTT
>2055371IH	GAGCAAGAGG	TGGCAGAGGA	GCGGAGAAAT	GCTCTCTTCC	CAGAGGTCTT
>3253860				GCTCTCTTCC	
>1753756				GCTCTCTTCC	
Consensus				GCTCTCTTCC	
>g2804590	CTCCCCAACG	CCAGATGAGA	ACTCTGACCA	GAACTCCAGG	AGCTCCTCCC
>2055371IH	CTCCCCAACG	CCAGATGAGA	ACTCTGACCA	GAACTCCAGG	AGCTCCTCCC
>3253860	С				
>1753756	CTCCCCAACG	CCAGATGAGA	ACTCTGACCA	GAACTCCAGG	AGCTCCTCCC
Consensus	CTCCCCAACG	CCAGATGAGA	ACTCTGACCA	GAACTCCAGG	AGCTCCTCCC

### Figure 1-E

>g2804590	AGGCATCCGG				
>g2804590 >g2804590		CATCACCCCC	AGTTACTCGG	ጥርጥርጥር እርጥር	ጥርርርጥጥርጥጥር
>2055371IH			AGTTACTCGG		
>1753756			AGTTA: TCGG		
>1887713	AGGCATCCGG	CATCACGGGC	AGITA. TCGG		TCCCTTCTTC
	N C C C N TI C C C C	CAMCACCCCC	AGTTACTCGG	_	
Consensus	AGGCATCCGG	CATCACGGGC	AGTIACTOGG	IGICIGAGIC	recerrence
>g2804590	AGCCCCATCC	ACCTACACTC	AAACGTGGCG	TGGACAGTGG	AAGATCCAGT
>2055371IH			AAACGTGGCG		
>1887713			AAACGTGGCG		
Consensus	AGCCCCATCC	ACCTACACTC	AAACGTGGCG	TGGACAGTGG	AAGATCCAGT
>g2804590 ·	GGACAGTGCT	CCTCCCGGGC	AGAGAAAGAA	GGAGCAATGG	
>g2804590				G	TACGCTGGCA
>2055371IH	GGACAGTGCT	CCTCCCGGGC	AGAGAAAGAA	GGAGCAATGG	TACGCTGGCA
>1887713	GGACAGTGCT	CCTCCCGGGC	AGAGAAAGAA	GGAGCAATGG	TACGCTGGCA
Consensus	GGACAGTGCT	CCTCCCGGGC	AGAGAAAGAA	GGAGCAATGG	TACGCTGGCA
>g2804590	TCAACCCCTC	GGACGGTATC	AACTCAGAGG	Т	
>g2804590			AGG	TCCTGGAAGC	CATACGGGTG
>2055371IH	TCAACCCCTC	GGACGGTATC	AACTCAGAGG	TCCTGGAAGC	CATACGGGTG
>1887713	TCAACCCCTC	GGACGGTATC	AACTCAGAGG	TCCTGGAAGC	CATACGGGTG
Consensus	TCAACCCCTC	GGACGGTATC	AACTCAGAGG	TCCTGGAAGC	CATACGGGTG
			•		
>g2804590	ACCCGTCACA	AGAACGCCAT	GGCAGAGCGC	TGGGAATCCC	GCATCTACGC
>2055371IH	ACCCGTCACA	AGAACGCCAT	GGCAGAGCGC	TGGGAATCCC	GCATCTACGC
>1887713	ACCCGTCACA	AGAACGCCAT	GGCAGAGCGC	TGGGAATCCC	GCATCTACGC
>1803052			C	TGGGAATCCC	GCATCTACGC
Consensus	ACCCGTCACA	AGAACGCCAT	GGCAGAGCGC	TGGGAATCCC	GCATCTACGC
>g2804590			CCTCGGGATG		•
>2055371IH			CCTCGGGATG		
>1887713			CCTCGGGATG		
>1803052			CCTCGGGATG		
Consensus	CAGTGAGGAG	GATGACTGAG	CCTCGGGATG	GGGCGCCCAC	CCCCTGCCCT
2004500	000000000000000000000000000000000000000	maamaaa	magaa a caaa	**************************************	CCCACCOTA
>g2804590			TGCCAAGACC TGCCAAGACC		
>2055371IH			TGCCAAGACC	ATCGCCAAGC	CCCCACCCTA
>1887713	GCCCTGACCC		magan Na Na	************	000000000000000000000000000000000000000
>1803052			TGCCAAGACC		
Consensus	GCCCTGACCC	TCGTGGGAAC	TGCCAAGACC	ATCGCCAAGC	CCCCACCCTA
>g2804590	GGAAATGGGT	CCTAGGTCCA	GGATCCAAGA	ACCACAGCTC	ATCTGCCAAC
>2055371IH					ATCTGCCAAC
>1803052					ATCTGCCAAC
Consensus					ATCTGCCAAC
001100110410	30.22110001	501501.00A			
>g2804590	AATCCCACCA	TGGGCACATT	TGGGACTGTT	GGGTTTTTCG	TTTCCGTTTC
>2055371IH	AATCCCACCA	TGGGCACATŤ	TGGGACTGTT	GGGTTTTTCG	TTTCCGTTTC
>1803052	AATCCCACCA	TGGGCACATT	TGGGACTGTT	GGGTTTTTCG	TTTCCGTTTC
Consensus	AATCCCACCA	TGGGCACATT	TGGGACTGTT	GGGTTTTTCG	TTTCCGTTTC

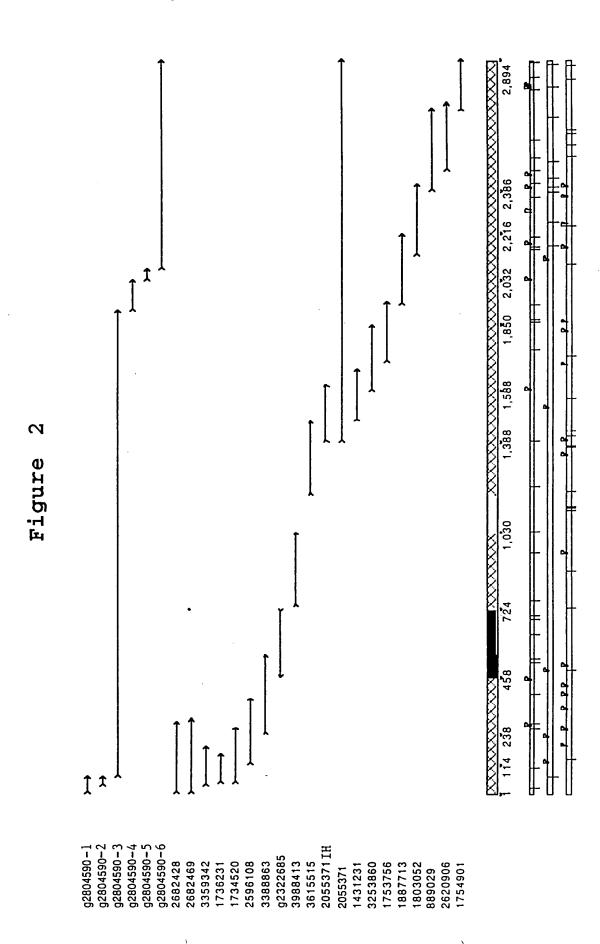
### Figure 1-F

>g2804590	TATCTTCCTT	TAGAAATGTT	TCTGCCTTTG	GGGTCTAAAG	CTTTTGGGGA
>2055371IH	TATCTTCCTT				
>1803052	TATCTTCCTT				
>889029					CTTTTGGGGA
Consensus	TATCTTCCTT	TAGAAATGTT	TCTGCCTTTG	GGGTCTAAAG	CTTTTGGGGA
Consensus					
>g2804590	TGAAATGGGA	CCCCTGCTGA	TTCTTTCTGC	TTCTAAGACT	TTGCCAAATG
>2055371IH	TGAAATGGGA				TTGCCAAATG
>1803052	TGAAATGGGA				
>889029	TGAAATGGGA		TTCTTTCTGC	TTCTAAGACT	TTGCCAAATG
Consensus			TTCTTTCTGC		
00040					
>g2804590	CCCTGGGTCT	AAGAAAGAAA	GAGACCCGCT	CCTCCACTTT	CAGGTGTAAT
>2055371IH	CCCTGGGTCT	AAGAAAGAAA	GAGACCCGCT	CCTCCACTTT	CAGGTGTAAT
>889029			GAGACCCGCT		CAGGTGTAAT
>2620906			GAGACCCGCT		CAGGTGTAAT
Consensus	CCCTGGGTCT		GAGACCCGCT		CAGGTGTAAT
Consensus	0001000101				
>g2804590	TTGCTTCCGC	TAGTCTGAGG	GCAGAGGGAC	CGGTCAAAGA	GGGTGGCACA
>2055371IH			GCAGAGGGAC		
>889029	TTGCTTCCGC		GCAGAGGGAC		
>2620906	TTGCTTCCGC		GCAGAGGGAC		
Consensus	TTGCTTCCGC		GCAGAGGGAC		
Consensus	1100110000	11101010101			
>g2804590	GATCGCAGCA	CCTTGAGGGG	CTGCGGGTCT	GAGGGAGGAG	ACACTCAGCT
>2055371IH					ACACTCAGCT
>889029					ACACTCAGCT
>2620906					ACACTCAGCT
Consensus					ACACTCAGCT
Conscinado	0				
>g2804590	CCTCCCTCTG	AGAAGTCCCA	AGCTGAGAGG	GGAGACCTGC	CCCTTTCCAA
>2055371IH					CCCTTTCCAA
>889029	CCTCCCTCTG	AGAAGTCCCA	AGCTGAGAGG	GGAGACCTGC	CCCTTTCCAA
>2620906	CCTCCCTCTG	AGAAGTCCCA	AGCTGAGAGG	GGAGACCTGC	CCCTTTCCAA
Consensus	CCTCCCTCTG	AGAAGTCCCA	AGCTGAGAGG	GGAGACCTGC	CCCTTTCCAA
>g2804590	CCCTGGGAAA	CCATCCAGTC			CCAGTGCTGG
>2055371IH	CCCTGGGAAA	CCATCCAGTC	TGAGGGAGGA	GGCCAAACTC	CCAGTGCTGG
>889029	CCCTGGGAAA	CCATCCAGTC	TGAGGGAGGA	GGCCAAACTT	CCAGTGCTGG
>2620906	CCCTGGGAAA	CCATCCAGTO	TGAGGGAGGA	GGCCAAACTC	CCAGTGCTGG
Consensus	CCCTGGGAAA	CCATCCAGTO	TGAGGGAGGA	GGCCAAACTC	CCAGTGCTGG
•					
>g2804590					CCAGCCACAC
>2055371IH	GGGTCCCTGT	GCAGCCCTCA	AACCCTTCAC	CTTGGTGCAC	CCAGCCACAC
>889029	GGGTCCCTGT	GCA			
>2620906	GGGTCCCTGT	GCAGCCCTCA	AACCCTTC		
>1754901	GGTCCCTGT	GCAGCCCTCA	AACCCTTCAC	CTTGGTGCAC	CCAGCCACAC
Consensus					CCAGCCACAC
50115011545					
>g2804590	CTGGTGGACA	CAAAGCTCTC	CACATCGATAC	GATCCCATGA	GGATGGTCCC
>2055371IH					GGATGGTCCC
>1754901					GGATGGTCCC
Consensus					GGATGGTCCC

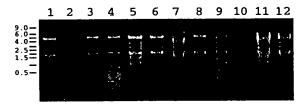
Reagents and Methods Useful for Detecting Diseases of the Gastrointestinal Tract 6065.US.D3 P.A. Billing-Medel, et al.

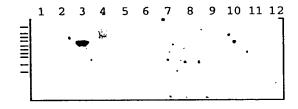
# Figure 1-G

>g2804590 >2055371IH >1754901 Consensus	CTTCACCTGG CTTCACCTGG	GAGAAAAGTG GAGAAAAGTG GAGAAAAGTG GAGAAAAGTG	ACCCAGTTTA ACCCAGTTTA	GGAGCTGGAG GGAGCTGGAG	GGGGGTCTTT GGGGGTCTTT
>g2804590 >2055371IH >1754901 Consensus	GTCCCCCACC GTCCCCCACC	CCCAAACTGC CCCAAACTGC CCCAAACTGC	CCTGAAATAA CCTGAAATAA	ACCTGGAGTG ACCTGGAGTG	AGCTGCCCA AGCTGCC



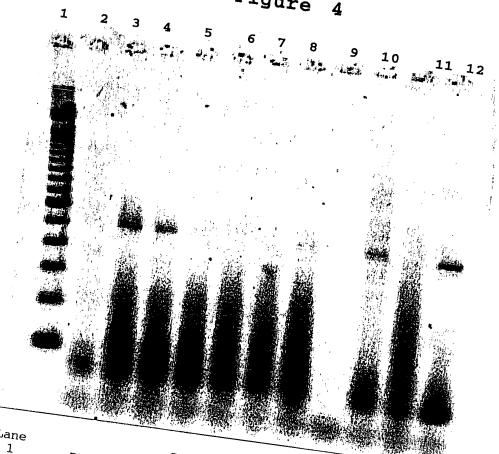
### Figure 3





Lane	Tissue
1	Bladder
2	Breast
3	Colon
4	Kidney
5	Liver
6	Lung
7	Muscle
8	Ovary
9	Placenta
10	Prostate
11	Spleen
12	Testis

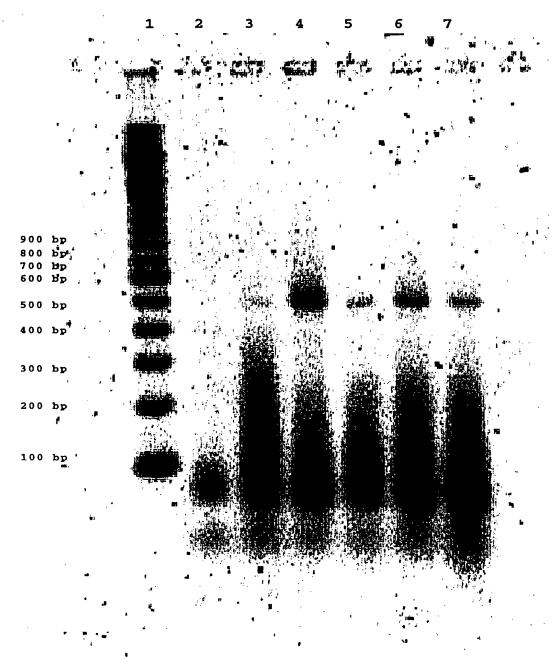
# Figure 4



Lane	The state of the s
1 2 3 4 5 6 7 8 9 10 11	Contents DNA Molecular Weight Marker Placental DNA control normal colon cancer colon cancer breast cancer breast BPH prostate cancer prostate BPH prostate normal lung normal lung cancer lung

Reagents and Methods Useful for Detecting
Diseases of the Gastrointestinal Tract
6065.US.D3
P.A. Billing-Medel, et al.

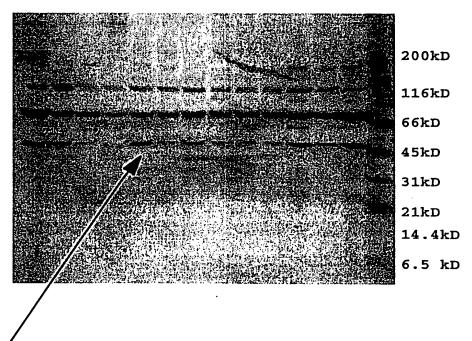
# Figure 5



Lane	Contents
1	DNA Molecular Weight Marker (100 bp)
2	placental DNA control
3	normal colon
4	cancer colon
5	cancer colon
6	normal colon
7	cancer colon

### Figure 6

#### 1 2 3 4 5 6 7 8 9 10 11 12 13 14



Lane	1	Endometrial cancer
Lane	2	Breast cancer
Lane	3	Normal lung
Lane	4	BPH prostate
Lane	5	Ovarian cancer
Lane	6	Normal bladder
Lane	7	Bladder cancer
Lane	8	Normal colon
Lane	9	Normal colon
Lane	10	Normal colon
Lane	11	Colon cancer
Lane	12	Colon cancer
Lane	13	Colon cancer
Lane	1 <i>1</i>	Markers